

From: Bryan J Pesta
To: John Fuerst
Subject: Re: Faculty access to an NIH database (dbGaP)
Date: Wednesday, April 11, 2018 12:51:50 PM
Attachments: image001.jpg



John,

Excellent for the strategy, and for noon on Tuesday.

Bryan

From: John Fuerst <j122177@hotmail.com>
Sent: Wednesday, April 11, 2018 12:48:16 PM
To: Bryan J Pesta
Subject: Re: Faculty access to an NIH database (dbGaP)

Tuesday at noon is fine with me.

As for the RUS, are you ok with my plan (according to which we look at the sex x g x brain size association and include population structure as a control and note the PC1 ancestry x IQ associations in the supplementary file)? This is a safe analysis. If so, I will write up a simple summary and note the variables needed. Otherwise, I can come up with something else.

John

From: Bryan J Pesta <b.pest@csuohio.edu>
Sent: Wednesday, April 11, 2018 11:40 AM
To: John Fuerst
Subject: Re: Faculty access to an NIH database (dbGaP)

This section might be important too. It was just above what I pasted in my prior email.

Bryan

From: Bryan J Pesta
Sent: Wednesday, April 11, 2018 4:40:33 AM
To: John Fuerst
Subject: Re: Faculty access to an NIH database (dbGaP)

Hi John,

How about Tuesday at noon?

I'm making good progress on the application. However, I'm stuck at this part:

Please enter your RUS in the area below. The RUS should be one or two paragraphs in length and include research objectives, the study design, and an analysis plan (including the phenotypic characteristics that will be tested for association with genetic variants). If you are requesting multiple datasets, please describe how you will use them. Examples of RUS can be found at [GDS website](#). Please limit your RUS to 2200 characters.

I am requesting permission to use cloud computing to carry out the research as described in my Research Use Statement.

Non-technical summary *What is a "non-technical summary of your Research Use Statement?"

Please enter below a non-technical summary of your RUS suitable for understanding by the general public (written at a high school reading level or below). Please limit your non-technical summary to 1100 characters.

It seems like your summary is for the RUS, but we need something toned down for the "non-technical" summary. Could you draft this?

Finally, in the RUS you attached, should the word "brain" appear in line 4 before the words "size/volume".

Thanks!

Bryan

From: John Fuerst <j122177@hotmail.com>
Sent: Tuesday, April 10, 2018 1:15:40 PM
To: Bryan J Pesta
Subject: Re: Faculty access to an NIH database (dbGaP)

Hi Bryan,

I am not sure if they will require one to be submitted. In case that they did, I proposed the attached.

If you are comfortable with publishing such a paper (and reporting the admixture results in a table in the supplementary file), then I say we go with it. If not, I can come up with something else.

Next Tuesday at Prosperity sounds fine. What time works for you?

Sincerely,
John

From: Bryan J Pesta <b.pest@csuohio.edu>
Sent: Tuesday, April 10, 2018 1:31 PM
To: John Fuerst
Subject: Re: Faculty access to an NIH database (dbGaP)

John,

You sent me a summary/abstract awhile ago. Should I be using that when I apply?

Also, what kind of food do you like or not like? Have you ever been to Prosperity (Starkweather and W. 11)? It has Polish and American food, plus a really nice patio, depending on the weather.

Next Tuesday (4/17) is open for me. Let me know-- I'm flexible.

From: John Fuerst <j122177@hotmail.com>
Sent: Monday, April 9, 2018 3:59:47 PM
To: Bryan J Pesta
Subject: Re: Faculty access to an NIH database (dbGaP)

Bryan,

Glad to here. We are at about 1.40 m here https://www.youtube.com/watch?v=m0xp_cCO7kA

You will have to select the survey you are applying for which is "Trajectories of complex phenotypes".

This does not require IRB approval, so we will be able to skip that step.

As for meeting up, ya sure, name a place and time.

Later,

John

From: Bryan J Pesta <b.pest@csuohio.edu>
Sent: Monday, April 9, 2018 3:24 PM
To: John Fuerst

Subject: Fw: Faculty access to an NIH database (dbGaP)

Hi John,

Making some progress -- see below.

What's the next step, and are you available next week for lunch (ideally Tremont, or Ohio City, but I understand if you don't want to go west)?

Bryan

From: Bryan J Pesta

Sent: Monday, April 9, 2018 9:35:42 AM

To: Lisa N Franklin

Subject: Re: Faculty access to an NIH database (dbGaP)

Thank you!

Bryan

From: Lisa N Franklin

Sent: Monday, April 9, 2018 9:17:27 AM

To: Bryan J Pesta

Subject: RE: Faculty access to an NIH database (dbGaP)

Good morning Bryan,

I've set you up with an eRA Commons account (NIH program), which should help you with accessing the data you are looking for. You will receive an email to complete the setup of your account. Let me know if you have any additional questions.

Thanks,

Lisa

Lisa Franklin, MBA

Assistant Director, Sponsored Programs & Research Services
2121 Euclid Avenue
Parker Hannifin Hall 220
Cleveland, OH 44115
216-687-3607
lfranklin@csuohio.edu



From: Mary Jane Karpinski

Sent: Monday, April 09, 2018 8:33 AM

To: Bryan J Pesta <b.pest@csuohio.edu>; Sponsored Programs and Research Services <spers@csuohio.edu>

Cc: Lisa N Franklin <l.franklin@csuohio.edu>

Subject: RE: Faculty access to an NIH database (dbGaP)

Good morning Bryan,

I am copying Lisa Franklin, Assistant Director of Sponsored Programs to help with your request.

Sincerely,
Mary Jane

Mary Jane Karpinski
Administrative Coordinator
Sponsored Programs & Research Services
Cleveland State University
Parker Hannifin Hall, 2nd floor

Phone: 216-687-3624/Fax: 216-687-9382

From: Bryan J Pesta
Sent: Sunday, April 08, 2018 11:15 AM
To: Sponsored Programs and Research Services <sprs@csuohio.edu>; Mary Jane Karpinski <m.karpinski2@csuohio.edu>
Subject: Faculty access to an NIH database (dbGaP)

Greetings,

My name is Bryan Pesta, and I am a professor here in the college of business.

I'm hoping you can help. I'm interested in accessing some NIH data-- specifically, certain data sets in the DBGAP (database of Genotypes and Phenotypes). I've never done this before, and the process seems complicated.

It seems like I must first contact and coordinate with CSU's "signing official" (SO), and then complete some paperwork. As a first step, can you tell me who our SO is, or provide any other information on how someone new to all this can go through the process of getting access to these data?

Thanks sincerely for your time,

Bryan Pesta

<https://www.ncbi.nlm.nih.gov/gap>

[Home - dbGaP - NCBI](https://www.ncbi.nlm.nih.gov/gap)

www.ncbi.nlm.nih.gov

The database of Genotypes and Phenotypes (dbGaP) was developed to archive and distribute the data and results from studies that have investigated the interaction of genotype and phenotype in Humans.